

Sequence Listing

Q1
<110> Lasky, Laurence A.
Dowbenko, Donald J.

<120> Tyrosine Phosphorylated Cleavage Furrow-Associated
Proteins (PSTPIPs)

<130> P1066P2

<140> US 09/068,377
<141> 1999-05-08

<150> US 08/938,300
<151> 1997-09-29

<150> US 08/798,419
<151> 1997-02-07

<160> 73

<210> 1
<211> 415
<212> PRT
<213> Mus Musculus

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Phe Thr Ala His Thr Gly Tyr Glu Val Leu Leu Gln Arg Leu Leu
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Asp Gly Arg Lys Met Cys Lys Asp Val Glu Glu Leu Leu Arg Gln
35 40 45
Arg Ala Gln Ala Glu Glu Arg Tyr Gly Lys Glu Leu Val Gln Ile
50 55 60
Ala Arg Lys Ala Gly Gly Gln Thr Glu Met Asn Ser Leu Arg Thr
65 70 75
Ser Phe Asp Ser Leu Lys Gln Gln Thr Glu Asn Val Gly Ser Ala
80 85 90
His Ile Gln Leu Ala Leu Ala Arg Glu Glu Leu Arg Ser Leu
95 100 105
Glu Glu Phe Arg Glu Arg Gln Lys Glu Gln Arg Lys Lys Tyr Glu
110 115 120
Ala Ile Met Asp Arg Val Gln Lys Ser Lys Leu Ser Leu Tyr Lys
125 130 135
Lys Thr Met Glu Ser Lys Lys Ala Tyr Asp Gln Lys Cys Arg Asp
140 145 150

Alf
cont.

Ala Asp Asp Ala Glu Gln Ala Phe Glu Arg Val Ser Ala Asn Gly
155 160 165

His Gln Lys Gln Val Glu Lys Ser Gln Asn Lys Ala Lys Gln Cys
170 175 180

Lys Glu Ser Ala Thr Glu Ala Glu Arg Val Tyr Arg Gln Asn Ile
185 190 195

Glu Gln Leu Glu Arg Ala Arg Thr Glu Trp Glu Gln Glu His Arg
200 205 210

Thr Thr Cys Glu Ala Phe Gln Leu Gln Glu Phe Asp Arg Leu Thr
215 220 225

Ile Leu Arg Asn Ala Leu Trp Val His Cys Asn Gln Leu Ser Met
230 235 240

Gln Cys Val Lys Asp Asp Glu Leu Tyr Glu Glu Val Arg Leu Thr
245 250 255

Leu Glu Gly Cys Asp Val Glu Gly Asp Ile Asn Gly Phe Ile Gln
260 265 270

Ser Lys Ser Thr Gly Arg Glu Pro Pro Ala Pro Val Pro Tyr Gln
275 280 285

Asn Tyr Tyr Asp Arg Glu Val Thr Pro Leu Ile Gly Ser Pro Ser
290 295 300

Ile Gln Pro Ser Cys Gly Val Ile Lys Arg Phe Ser Gly Leu Leu
305 310 315

His Gly Ser Pro Lys Thr Thr Pro Ser Ala Pro Ala Ala Ser Thr
320 325 330

Glu Thr Leu Thr Pro Thr Pro Glu Arg Asn Glu Leu Val Tyr Ala
335 340 345

Ser Ile Glu Val Gln Ala Thr Gln Gly Asn Leu Asn Ser Ser Ala
350 355 360

Gln Asp Tyr Arg Ala Leu Tyr Asp Tyr Thr Ala Gln Asn Ser Asp
365 370 375

Glu Leu Asp Ile Ser Ala Gly Asp Ile Leu Ala Val Ile Leu Glu
380 385 390

Gly Glu Asp Gly Trp Trp Thr Val Glu Arg Asn Gly Gln Arg Gly
395 400 405

Phe Val Pro Gly Ser Tyr Leu Glu Lys Leu
410 415

<210> 2
<211> 2100
<212> DNA
<213> Mus Musculus

All
Done

<400> 2

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gaacctcata acaactcaaa caaattctca agcgcttca caaccaattg 200
cctcctctaa cgttcatgat aacttcatga ataatgaaat cacggctagt 250
aaaattgatg atggtataa ttcaaaaacca ctgtcacctg gttggacgga 300
ccaaactgct tataacgcgt ttggaatcac tacagggatg tttatcacca 350
ctacaatgga tcatgtat aactatctat tcgatgatga agataccca 400
ccaaacccaa aaaaagaggg tgggtcgacc cacgcgtccg gtccttcct 450
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caagaaggca tatgaccaga agtgcaggga tgcagatgat gctgagcagg 1150
ccttcgagcg tgtgagtgcc aatggccacc agaagcaagt agaaaagagc 1200
cagaacaaag ccaagcagtg caaggagtca gccacagagg cagaaagagt 1250
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aggagcaccc gactacctgt gaggccttcc agttgcagga gtttgaccgg 1350
ctcaccatcc tccgcaatgc cctgtgggtg cactgtaacc agctctccat 1400
gcagtgtgtc aaggatgatg agctctatga ggaagtgcgg ctgacccttg 1450

Ch
Cont
agggctgtga tgtggaaggt gacatcaatg gcttcatcca gtccaaagac 1500
actggcagag agcccccagc tccgggtgcct tatcagaact actatgacag 1550
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tgataaaagag gttctctggg ctgctacatg gaagtcccaa gaccacac 1650
tctgctccctg ctgcttccac agagactctg actcccaccc ctgagccgaa 1700
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tctgatgagc tggacatttc cgcgggagac atcctggcgg tcattcctgga 1850
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tccctgggtc gtacttggag aagctctgag gaaaggctag cagtctccac 1950
atacctccgc cctgactgtg aggtcaggac tggcctttc catcaccgac 2000
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gtgctggcta ctctcaataa atgtctccca gaaggaaaaaa aaaaaaaaaa 2100

<210> 3
<211> 48
<212> PRT
<213> Mus Musculus

<400> 3
Leu Tyr Asp Tyr Thr Ala Gln Asn Ser Asp Glu Leu Asp Ile Ser
1 5 10 15
Ala Gly Asp Ile Leu Ala Val Ile Leu Glu Gly Glu Asp Gly Trp
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Trp Thr Val Glu Arg Asn Gly Gln Arg Gly Phe Val Pro Gly Ser
35 40 45
Tyr Leu Arg
48

<210> 4
<211> 50
<212> PRT
<213> Homo sapien

<400> 4
Leu Tyr Gln Tyr Ile Gly Gln Asp Val Asp Glu Leu Ser Phe Asn
1 5 10 15
Val Asn Glu Val Ile Glu Ile Leu Ile Glu Asp Ser Ser Gly Trp
20 25 30
Trp Lys Gly Arg Leu His Gly Gln Glu Gly Leu Phe Pro Gly Asn
35 40 45

Al
Conn

Tyr Val Glu Lys Ile
50

<210> 5
<211> 50
<212> PRT
<213> Homo sapien

<400> 5
Leu Tyr Asp Tyr Gln Glu Lys Ser Pro Arg Glu Val Thr Met Lys
1 5 10 15

Lys Gly Asp Ile Leu Thr Leu Leu Asn Ser Thr Asn Lys Asp Trp
20 25 30

Trp Lys Val Glu Val Asn Asp Arg Gln Gly Phe Val Pro Ala Ala
35 40 45

Tyr Val Lys Lys Leu
50

<210> 6
<211> 50
<212> PRT
<213> Homo sapien

<400> 6
Leu Tyr Asp Tyr Gln Gly Glu Gly Ser Asp Glu Leu Ser Phe Asp
1 5 10 15

Pro Asp Asp Ile Ile Thr Asp Ile Glu Met Val Asp Glu Gly Trp
20 25 30

Trp Arg Gly Gln Cys Arg Gly His Phe Gly Leu Phe Pro Ala Asn
35 40 45

Tyr Val Lys Leu Leu
50

<210> 7
<211> 48
<212> PRT
<213> Homo sapien

<400> 7
Leu Tyr Asp Tyr Gln Ala Ala Gly Asp Asp Glu Ile Ser Phe Asp
1 5 10 15

Pro Asp Asp Ile Ile Thr Asn Ile Glu Met Ile Asp Asp Gly Trp
20 25 30

Trp Arg Gly Val Cys Lys Gly Arg Tyr Gly Leu Phe Pro Ala Asn
35 40 45

Tyr Val Glu
48

Alb
Onc

<210> 8
<211> 8
<212> PRT
<213> Artificial Sequence

<220>
<221> Artificial Sequence
<222> 1-8
<223> Amino acid epitope tag

<400> 8
Asp Tyr Lys Asp Asp Asp Asp Lys
1 5 8

<210> 9
<211> 33
<212> DNA
<213> Artificial Sequence

<220>
<221> Artificial Sequence
<222> 1-33
<223> Synthetic oligonucleotide probe

<400> 9
cgcggatcca ccatgatggc ccagctgcag ttc 33

<210> 10
<211> 45
<212> DNA
<213> Artificial Sequence

<220>
<221> Artificial Sequence
<222> 1-45
<223> Synthetic oligonucleotide probe

<400> 10
gtacgcgtcg actcaacttgt catcgatcg cttgtatcg agctt 45

<210> 11
<211> 18
<212> DNA
<213> Artificial Sequence

<220>
<221> Artificial Sequence
<222> 1-18
<223> Synthetic oligonucleotide probe

<400> 11
tgcctttctc tccacagg 18

<210> 12
<211> 36
<212> DNA
<213> Artificial Sequence

<220>
<221> Artificial Sequence
<222> 1-36
<223> Synthetic oligonucleotide probe

<400> 12
ctccttgagg ttctactagt gggggctgg 36

<210> 13
<211> 39
<212> DNA
<213> Artificial Sequence

<220>
<221> Artificial Sequence
<222> 1-39
<223> Synthetic oligonucleotide probe

<400> 13
gcggccgcac tagtatccag tctgtgctcc atctgttac 39

<210> 14
<211> 17
<212> DNA
<213> Artificial Sequence

<220>
<221> Artificial Sequence
<222> 1-17
<223> Synthetic oligonucleotide probe

<400> 14
gcgtttggaa tcactac 17

<210> 15
<211> 41
<212> DNA
<213> Artificial Sequence

<220>
<221> Artificial Sequence
<222> 1-41
<223> Synthetic oligonucleotide probe

<400> 15
ttatagtttgcggccgctc accggtagtc ctgggctgat g 41

<210> 16
<211> 37
<212> DNA
<213> Artificial Sequence

<220>
<221> Artificial Sequence
<222> 1-37
<223> Synthetic oligonucleotide probe

<400> 16
gtacgcgtcg accgcactct acgactacac tgcacag 37

<210> 17
<211> 17
<212> DNA
<213> Artificial Sequence

<220>
<221> Artificial Sequence
<222> 1-17
<223> Synthetic oligonucleotide probe

<400> 17
ctctggcgaa gaagtcc 17

<210> 18
<211> 32
<212> DNA
<213> Artificial Sequence

<220>
<221> Artificial Sequence
<222> 1-32
<223> Synthetic oligonucleotide probe

<400> 18
gatcgaattc ccagaacctc aaggagaact gc 32

<210> 19
<211> 38
<212> DNA
<213> Artificial Sequence

<220>
<221> Artificial Sequence
<222> 1-38
<223> Synthetic oligonucleotide probe

<400> 19
gatcctcgag ttacacccgt gtccactctg ctggagga 38

<210> 20
<211> 20
<212> PRT
<213> Artificial Sequence

<220>
<221> Artificial Sequence
<222> 1-20
<223> Synthetic oligopeptide

<400> 20
Gly Phe Asn Leu Arg Ile Gly Arg Pro Lys Gly Pro Arg Asp Pro
1 5 10 15

Pro Ala Glu Trp Thr
20

<210> 21
<211> 19
<212> PRT
<213> Artificial Sequence

<220>
<221> Artificial Sequence
<222> 1-19
<223> Synthetic oligopeptide

<400> 21
Gly Phe Gly Asn Arg Phe Ser Lys Pro Lys Gly Pro Arg Asn Pro
1 5 10 15

Pro Ser Ala Trp
19

<210> 22
<211> 20
<212> PRT
<213> Artificial Sequence

<220>
<221> Artificial Sequence
<222> 1-20
<223> Synthetic oligopeptide

<400> 22
Gly Phe Gly Asn Arg Cys Gly Lys Pro Lys Gly Pro Arg Asp Pro
1 5 10 15

Pro Ser Glu Trp Thr
20

<210> 23
<211> 20
<212> PRT
<213> Artificial Sequence

<220>
<221> Artificial Sequence
<222> 1-20
<223> Synthetic oligopeptide

<400> 23
Gly Gly Val Leu Arg Ser Ile Ser Val Pro Ala Pro Pro Thr Leu
1 5 10 15

Pro Met Ala Asp Thr
20

<210> 24
<211> 36
<212> DNA
<213> Artificial Sequence

<220>
<221> Artificial Sequence

<222> 1-36
<223> Synthetic oligonucleotide probe

<400> 24
gtatatgtcc tggccagccc atggggttcc cagcag 36

<210> 25
<211> 36
<212> DNA
<213> Artificial Sequence

<220>
<221> Artificial Sequence
<222> 1-36
<223> Synthetic oligonucleotide probe

<400> 25
gcaggtcgac tctagattac acccgtgtcc actctg 36

<210> 26
<211> 907
<212> PRT
<213> *Saccharomyces Pombe*

<400> 26

Met	Leu	Thr	Lys	Ser	Leu	Gln	Gly	Ser	Glu	Asp	Ala	Gly	Met	Asp
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		20						25				30		
Ile	Asp	Glu	Phe	Tyr	Ala	Lys	Arg	Ala	Ser	Ile	Glu	Arg	Glu	Tyr
			35					40				45		
Ala	Ser	Lys	Leu	Gln	Glu	Leu	Ala	Ala	Ser	Ser	Ala	Asp	Ile	Pro
			50					55				60		
Glu	Val	Gly	Ser	Thr	Leu	Asn	Asn	Ile	Leu	Ser	Met	Arg	Thr	Glu
			65					70				75		
Thr	Gly	Ser	Met	Ala	Lys	Ala	His	Glu	Glu	Val	Ser	Gln	Gln	Ile
			80					85				90		
Asn	Thr	Glu	Leu	Arg	Asn	Lys	Ile	Arg	Glu	Tyr	Ile	Asp	Gln	Thr
			95					100				105		
Glu	Gln	Gln	Lys	Val	Val	Ala	Ala	Asn	Ala	Ile	Glu	Glu	Leu	Tyr
			110					115				120		
Gln	Lys	Lys	Thr	Ala	Leu	Glu	Ile	Asp	Leu	Ser	Glu	Lys	Lys	Asp
			125					130				135		
Ala	Tyr	Glu	Tyr	Ser	Cys	Asn	Lys	Leu	Asn	Ser	Tyr	Met	Arg	Gln
			140					145				150		
Thr	Lys	Lys	Met	Thr	Gly	Arg	Glu	Leu	Asp	Lys	Tyr	Asn	Leu	Lys
			155					160				165		

Ile	Arg	Gln	Ala	Ala	Leu	Ala	Val	Lys	Lys	Met	Asp	Ala	Glu	Tyr
170								175					180	
Arg	Glu	Thr	Asn	Glu	Leu	Leu	Leu	Thr	Val	Thr	Arg	Glu	Trp	Ile
185								190					195	
Asp	Arg	Trp	Thr	Glu	Val	Cys	Asp	Ala	Phe	Gln	His	Ile	Glu	Glu
200								205					210	
Tyr	Arg	Leu	Glu	Phe	Leu	Lys	Thr	Asn	Met	Trp	Ala	Tyr	Ala	Asn
215								220					225	
Ile	Ile	Ser	Thr	Ala	Cys	Val	Lys	Asp	Asp	Glu	Ser	Cys	Glu	Lys
230								235					240	
Ile	Arg	Leu	Thr	Leu	Glu	Asn	Thr	Asn	Ile	Asp	Glu	Asp	Ile	Thr
245								250					255	
Gln	Met	Ile	Gln	Asn	Glu	Gly	Thr	Gly	Thr	Thr	Ile	Pro	Pro	Leu
260								265					270	
Pro	Glu	Phe	Asn	Asp	Tyr	Phe	Lys	Glu	Asn	Gly	Leu	Asn	Tyr	Asp
275								280					285	
Ile	Asp	Gln	Leu	Ile	Ser	Lys	Ala	Pro	Ser	Tyr	Pro	Tyr	Ser	Ser
290								295					300	
Ser	Arg	Pro	Ser	Ala	Ser	Ala	Ser	Leu	Ala	Ser	Ser	Pro	Thr	Arg
305								310					315	
Ser	Ala	Phe	Arg	Pro	Lys	Thr	Ser	Glu	Thr	Val	Ser	Ser	Glu	Val
320								325					330	
Val	Ser	Ser	Pro	Pro	Thr	Ser	Pro	Leu	His	Ser	Pro	Val	Lys	Pro
335								340					345	
Val	Ser	Asn	Glu	Gln	Val	Glu	Gln	Val	Thr	Glu	Val	Glu	Leu	Ser
350								355					360	
Ile	Pro	Val	Pro	Ser	Ile	Gln	Glu	Ala	Glu	Ser	Gln	Lys	Pro	Val
365								370					375	
Leu	Thr	Gly	Ser	Ser	Met	Arg	Arg	Pro	Ser	Val	Thr	Ser	Pro	Thr
380								385					390	
Phe	Glu	Val	Ala	Ala	Arg	Pro	Leu	Thr	Ser	Met	Asp	Val	Arg	Ser
395								400					405	
Ser	His	Asn	Ala	Glu	Thr	Glu	Val	Gln	Ala	Ile	Pro	Ala	Ala	Thr
410								415					420	
Asp	Ile	Ser	Pro	Glu	Val	Lys	Glu	Gly	Lys	Asn	Ser	Glu	Asn	Ala
425								430					435	
Ile	Thr	Lys	Asp	Asn	Asp	Asp	Ile	Ile	Leu	Ser	Ser	Gln	Leu	Gln
440								445					450	

Pro Thr Ala Thr Gly Ser Arg Ser Ser Arg Leu Ser Phe Ser Arg
 455 460 465
 His Gly His Gly Ser Gln Thr Ser Leu Gly Ser Ile Lys Arg Lys
 470 475 480
 Ser Ile Met Glu Arg Met Gly Arg Pro Thr Ser Pro Phe Met Gly
 485 490 495
 Ser Ser Phe Ser Asn Met Gly Ser Arg Ser Thr Ser Pro Thr Lys
 500 505 510
 Glu Gly Phe Ala Ser Asn Gln His Ala Thr Gly Ala Ser Val Gln
 515 520 525
 Ser Asp Glu Leu Glu Asp Ile Asp Pro Arg Ala Asn Val Val Leu
 530 535 540
 Asn Val Gly Pro Asn Met Leu Ser Val Gly Glu Ala Pro Val Glu
 545 550 555
 Ser Thr Ser Lys Glu Glu Asp Lys Asp Val Pro Asp Pro Ile Ala
 560 565 570
 Asn Ala Met Ala Glu Leu Ser Ser Ser Met Arg Arg Arg Gln Ser
 575 580 585
 Thr Ser Val Asp Asp Glu Ala Pro Val Ser Leu Ser Lys Thr Ser
 590 595 600
 Ser Ser Thr Arg Leu Asn Gly Leu Gly Tyr His Ser Arg Asn Thr
 605 610 615
 Ser Ile Ala Ser Asp Ile Asp Gly Val Pro Lys Lys Ser Thr Leu
 620 625 630
 Gly Ala Pro Pro Ala Ala His Thr Ser Ala Gln Met Gln Arg Met
 635 640 645
 Ser Asn Ser Phe Ala Ser Gln Thr Lys Gln Val Phé Gly Glu Gln
 650 655 660
 Arg Thr Glu Asn Ser Ala Arg Glu Ser Leu Arg His Ser Arg Ser
 665 670 675
 Asn Met Ser Arg Ser Pro Ser Pro Met Leu Ser Arg Arg Ser Ser
 680 685 690
 Thr Leu Arg Pro Ser Phe Glu Arg Ser Ala Ser Ser Leu Ser Val
 695 700 705
 Arg Gln Ser Asp Val Val Ser Pro Ala Pro Ser Thr Arg Ala Arg
 710 715 720
 Gly Gln Ser Val Ser Gly Gln Gln Arg Pro Ser Ser Ser Met Ser
 725 730 735

Leu	Tyr	Gly	Glu	Tyr	Asn	Lys	Ser	Gln	Pro	Gln	Leu	Ser	Met	Gln
740														750
Arg	Ser	Val	Ser	Pro	Asn	Pro	Leu	Gly	Pro	Asn	Arg	Arg	Ser	Ser
755														765
Ser	Val	Leu	Gln	Ser	Gln	Lys	Ser	Thr	Ser	Ser	Asn	Thr	Ser	Asn
770														780
Arg	Asn	Asn	Gly	Gly	Tyr	Ser	Gly	Ser	Arg	Pro	Ser	Ser	Glu	Met
785														795
Gly	His	Arg	Tyr	Gly	Ser	Met	Ser	Gly	Arg	Ser	Met	Arg	Gln	Val
800														810
Ser	Gln	Arg	Ser	Thr	Ser	Arg	Ala	Arg	Ser	Pro	Glu	Pro	Thr	Asn
815														825
Arg	Asn	Ser	Val	Gln	Ser	Lys	Asn	Val	Asp	Pro	Arg	Ala	Thr	Phe
830														840
Thr	Ala	Glu	Gly	Glu	Pro	Ile	Leu	Gly	Tyr	Val	Ile	Ala	Leu	Tyr
845														855
Asp	Tyr	Gln	Ala	Gln	Ile	Pro	Glu	Glu	Ile	Ser	Phe	Gln	Lys	Gly
860														870
Asp	Thr	Leu	Met	Val	Leu	Arg	Thr	Gln	Glu	Asp	Gly	Trp	Trp	Asp
875														885
Gly	Glu	Ile	Ile	Asn	Val	Pro	Asn	Ser	Lys	Arg	Gly	Leu	Phe	Pro
890														900
Ser	Asn	Phe	Val	Gln	Thr	Val								
905														

<210> 27
 <211> 4
 <212> PRT
 <213> Artificial Sequence

<220>
 <221> Any amino acid
 <222> 2-3
 <223> Any amino acid

<400> 27
 Pro Xaa Xaa Pro
 1 4

<210> 28
 <211> 1613
 <212> DNA
 <213> Homo sapien

<400> 28
 acgatcacta tagggcgaat tgggcctcta gatgcgtgcgt cgagcggccg 50

ccagtgtgat ggatatctgc agaattcggc ttccatccta atacgactca 100
ctatagggct cgagcggccg cccgggcagg tctagaattc agcggccgct 150
gaattctctt tttcctcccc tcagaagctc ctctctggct cgtggctgcc 200
ttctgagtgt tgcatcggc gcccggccgg aaggggggcc tggccagcc 250
ctgccaggac tgggacgctg ctgctgacgc ctggccctcc atcaggccag 300
cctgtggcag gagagtgagc tttgcccgg cagacgcctg aggatgatgc 350
cccagctgca gttcaaagat gccttttgt gcagggactt cacagccac 400
acgggctacg aggtgctgct gcagcggctt ctggatggca ggaagatgtg 450
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acgggaagga gctggcgcag atgcacgga aggaggtgg ccagacggag 550
atcaactccc tgagggcctc ctttgactcc ttgaagcagc aaatggagaa 600
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ggagtctcga ggagttcgt gagaggcaga aggagcagag gaagaaggc 700
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acaagaaggc catggagtcc aagaagacat acgagcagaa gtgccgggac 850
gcggacgacg cggagcaggc ctgcggcgc attagcgcca acggccacca 900
gaagcaggtg gagaagagtc agaacaaggc caggcagtgc aaggactcgg 950
ccaccgaggc agagcgggta tacaggcaga gcattgcgc gctggagaag 1000
gtccgggctg agtgggagca ggagcaccgg accacctgtg aggcctttca 1050
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acagtaacca gctctccatg cagtgtgtca aggatgatga gctctacgag 1150
gaagtgcggc tgacgctgga aggctgcagc atagacgccc acatcgacag 1200
tttcatccag gccaagagca cgggcacaga gccccccagg ttctctggac 1250
tgctgcacgg aagtcccaag accacttcgt cagttctgc tggctccaca 1300
gagaccctga cccccacccc cgagcggaat gaggtgtct acacagccat 1350
cgcagtgcag gagatacagg gaaacccggc ctcaccagcc caggactacc 1400
ggcgctcta cgattataca gcgcagaacc cagatgagct ggacctgtcc 1450
gcgggagaca tccttggagg ggaggatggc tggctggactg tggagagggaa 1500

cgggcagcgt ggcttcgtcc ctgggttccta cctggagaag ctttgaggga 1550
aggccaggag ccccttcgga cctccgcctt gccagtggag ccagcagtgc 1600
ccccagcact gtc 1613

<210> 29
<211> 400
<212> PRT
<213> Homo sapien

<400> 29
Met Met Pro Gln Leu Gln Phe Lys Asp Ala Phe Trp Cys Arg Asp
1 5 10 15
Phe Thr Ala His Thr Gly Tyr Glu Val Leu Leu Gln Arg Leu Leu
20 25 30
Asp Gly Arg Lys Met Cys Lys Asp Met Glu Glu Leu Leu Arg Gln
35 40 45
Arg Ala Gln Ala Glu Glu Arg Tyr Gly Lys Glu Leu Val Gln Ile
50 55 60
Ala Arg Lys Ala Gly Gly Gln Thr Glu Ile Asn Ser Leu Arg Ala
65 70 75
Ser Phe Asp Ser Leu Lys Gln Gln Met Glu Asn Val Gly Ser Ser
80 85 90
His Ile Gln Leu Ala Leu Thr Leu Arg Glu Glu Leu Arg Ser Leu
95 100 105
Glu Glu Phe Arg Glu Arg Gln Lys Glu Gln Arg Lys Lys Gly Met
110 115 120
Ala Val Pro Arg Gln Ser Asp Cys Met Glu Val Lys Ser Pro Ser
125 130 135
Trp Glu Tyr Glu Ala Val Met Asp Arg Val Gln Lys Ser Lys Leu
140 145 150
Ser Leu Tyr Lys Lys Ala Met Glu Ser Lys Lys Thr Tyr Glu Gln
155 160 165
Lys Cys Arg Asp Ala Asp Asp Ala Glu Gln Ala Phe Glu Arg Ile
170 175 180
Ser Ala Asn Gly His Gln Lys Gln Val Glu Lys Ser Gln Asn Lys
185 190 195
Ala Arg Gln Cys Lys Asp Ser Ala Thr Glu Ala Glu Arg Val Tyr
200 205 210
Arg Gln Ser Ile Ala Gln Leu Glu Lys Val Arg Ala Glu Trp Glu
215 220 225

Gln	Glu	His	Arg	Thr	Thr	Cys	Glu	Ala	Phe	Gln	Leu	Gln	Glu	Phe
230							235						240	
Asp	Arg	Leu	Thr	Ile	Leu	Arg	Asn	Ala	Leu	Trp	Val	His	Ser	Asn
245							250						255	
Gln	Leu	Ser	Met	Gln	Cys	Val	Lys	Asp	Asp	Glu	Leu	Tyr	Glu	Glu
260							265						270	
Val	Arg	Leu	Thr	Leu	Glu	Gly	Cys	Ser	Ile	Asp	Ala	Asp	Ile	Asp
275							280						285	
Ser	Phe	Ile	Gln	Ala	Lys	Ser	Thr	Gly	Thr	Glu	Pro	Pro	Arg	Phe
290							295						300	
Ser	Gly	Leu	Leu	His	Gly	Ser	Pro	Lys	Thr	Thr	Ser	Ser	Ala	Ser
305							310						315	
Ala	Gly	Ser	Thr	Glu	Thr	Leu	Thr	Pro	Thr	Pro	Glu	Arg	Asn	Glu
320							325						330	
Gly	Val	Tyr	Thr	Ala	Ile	Ala	Val	Gln	Glu	Ile	Gln	Gly	Asn	Pro
335							340						345	
Ala	Ser	Pro	Ala	Gln	Asp	Tyr	Arg	Ala	Leu	Tyr	Asp	Tyr	Thr	Ala
350							355						360	
Gln	Asn	Pro	Asp	Glu	Leu	Asp	Leu	Ser	Ala	Gly	Asp	Ile	Leu	Glu
365							370						375	
Gly	Glu	Asp	Gly	Trp	Trp	Thr	Val	Glu	Arg	Asn	Gly	Gln	Arg	Gly
380							385						390	
Phe	Val	Pro	Gly	Ser	Tyr	Leu	Glu	Lys	Leu					
							395						400	

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Done*

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Artificial
Oligonucleotide

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